

# Scoring Biological Integrity

## *the California Stream Condition Index (CSCI)*



- **Scoring Tool Enhancements**
  - Update to O/E component
  - Integrating predictive MMI techniques
  - Our recommendations
- **Setting Thresholds**
- **Statewide and Regional Extent Estimates**
- **Questions for the panel**



# Technical Team

**\*Andy Rehn, *DFG-ABL***

**\*Raphael Mazor, *SCCWRP +DFG-ABL***

**Larry Brown, *USGS***

**Jason May, *USGS***

**David Herbst, *SNARL***

**Peter Ode, *DFG-WPCL/ABL***

**Ken Schiff, *SCCWRP***

**David Gillett, *SCCWRP***

**Eric Stein, *SCCWRP***

**Betty Fetscher, *SCCWRP***

**Kevin Lunde, *SF Water Board***

**\*\*\* *Scientific Review Panel***



How do we convert a list of species  
into a condition score?

NABS ([www.benthos.org](http://www.benthos.org))

# Qualities of a good scoring tool

## Technical Qualities

- precise
- accurate
- responsive

## Regulatory Qualities

- universally applicable
- easy to relate to ecological condition
- easy to compare to a standard

# **Two common approaches for quantifying biotic condition**

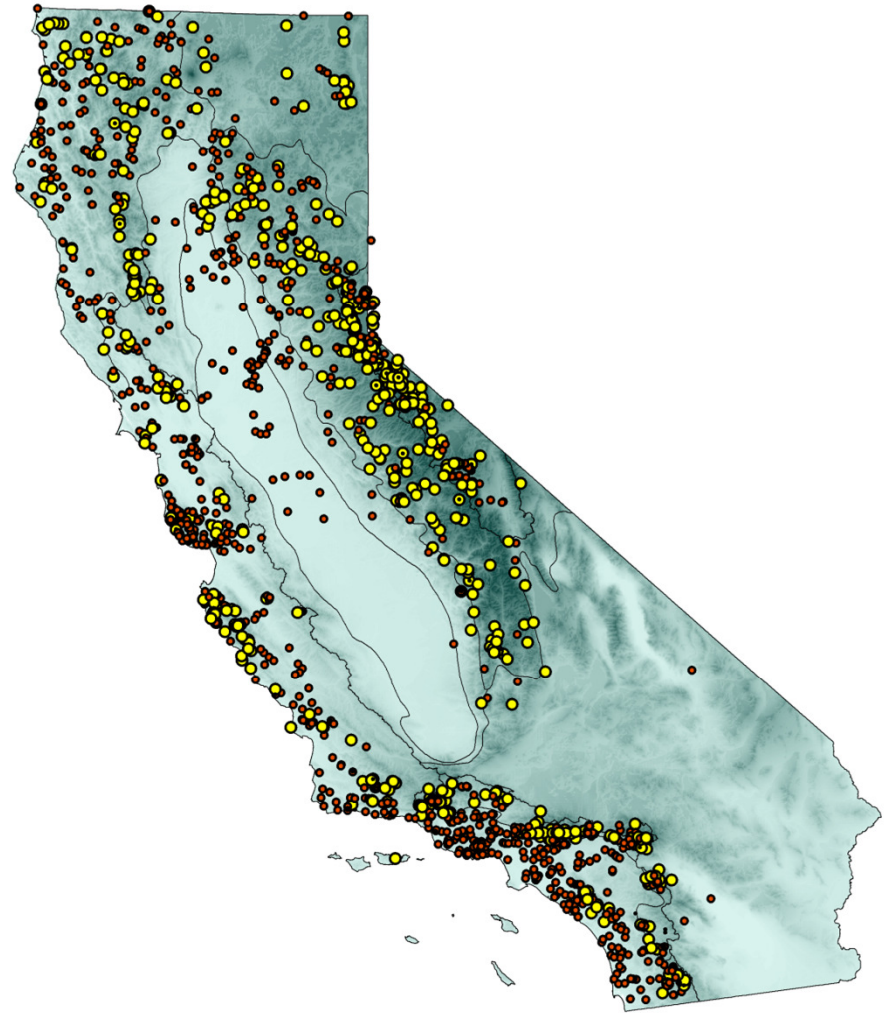
**Species loss indices** (e.g., O/E indices)

**Ecological structure indices** (e.g., multi-metric indices including IBIs)

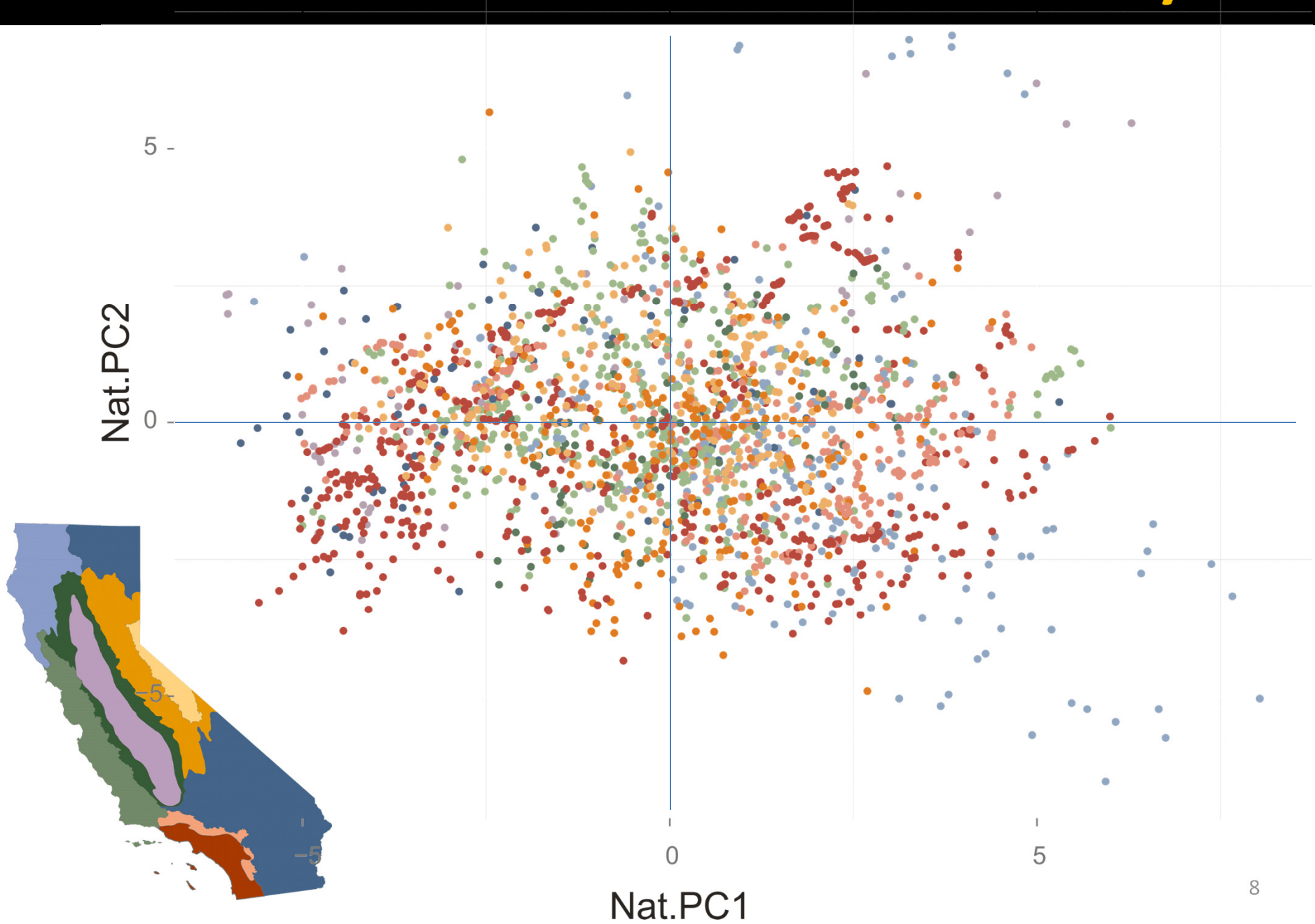


# Scoring tools rely on reference sites to establish expected conditions

- 485 reference sites used to develop scoring models
- Excellent coverage of CA's natural stream diversity

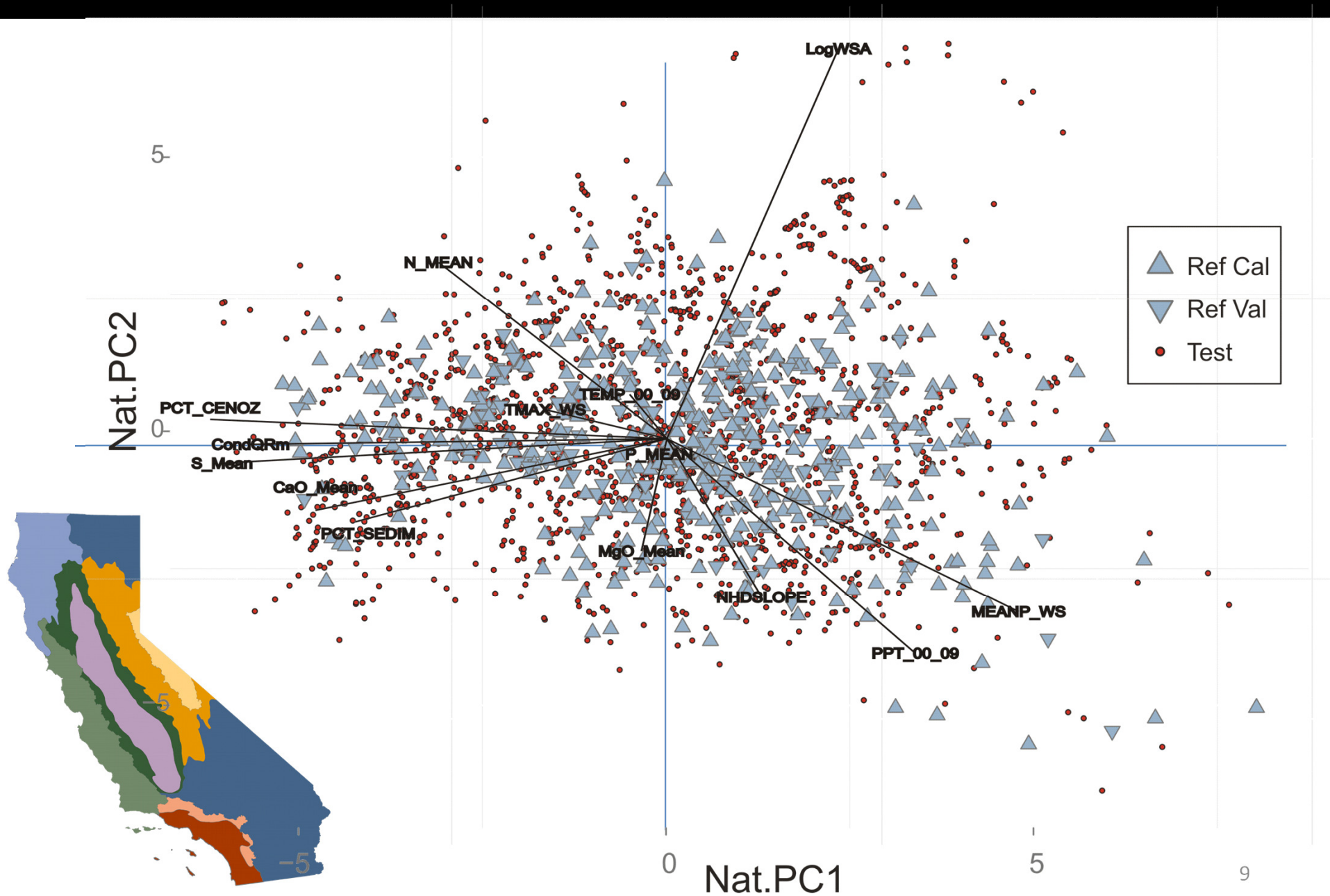


# Multivariate view of natural diversity





# Multivariate view of natural diversity



# Species Loss Index (O/E)

Compare number of **observed** (“O”) taxa to  
number of **expected** (“E”) taxa

**Step 1.** Cluster reference sites based on biological similarity

**Step 2.** Identify natural gradients that best explain clusters (=predictors)

**Step 3.** Use predictor values at test sites to predict species expected to be observed

*Index score is an estimate of taxonomic loss*

# O/E Update

- April index performed well
- **Reference pool adjustments:**
  - added sites to target under-represented gradients
  - dropped sites based on stakeholder feedback
- New discriminant functions model was not as precise as the April model
- Experimented with climatic sub-models, random forest techniques, predictor selection



# O/E Update

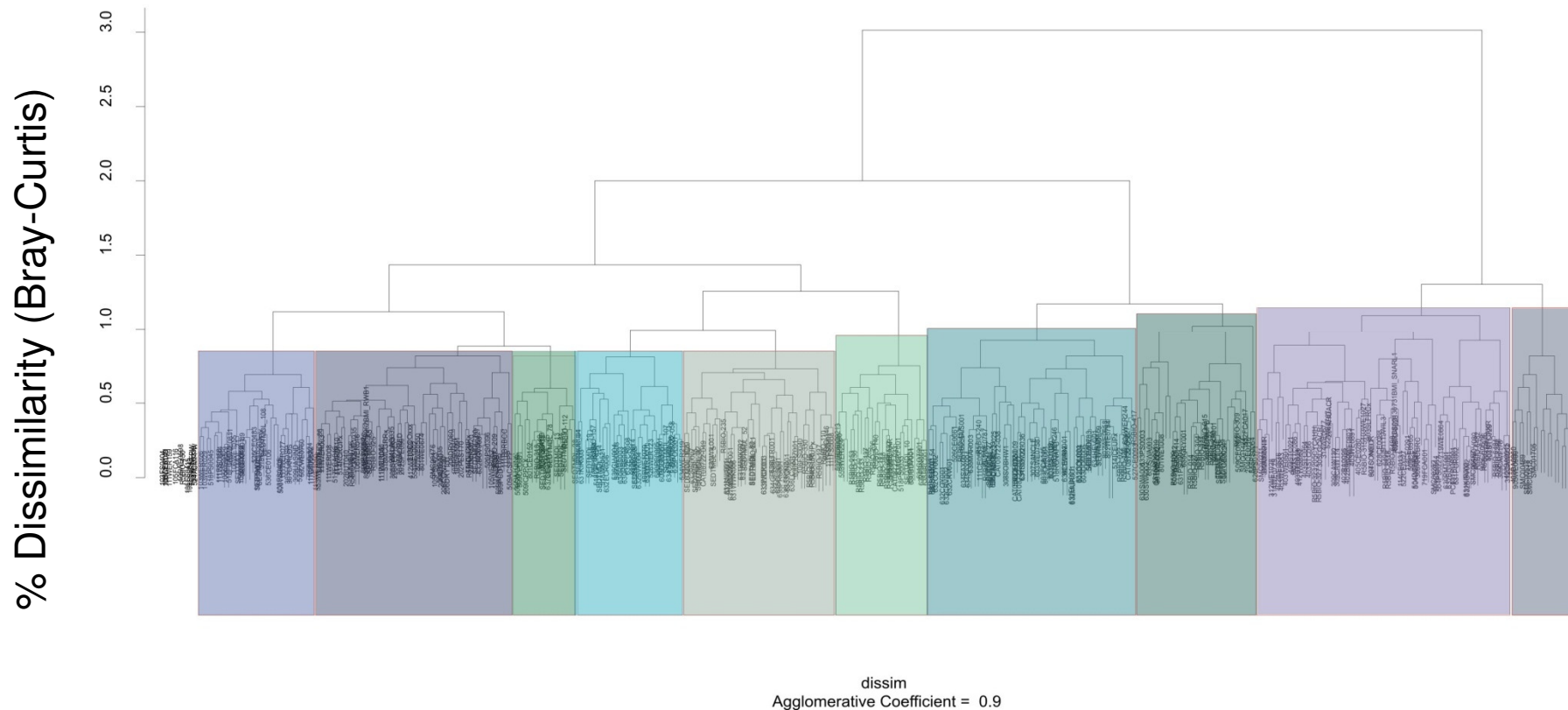
## Final Model (Random Forests, 10 clusters, 4 predictors):

- Average Monthly Temperature (2000-2009)
- Average Monthly Precipitation (2000-2009)
- Log Watershed Area
- Site Elevation

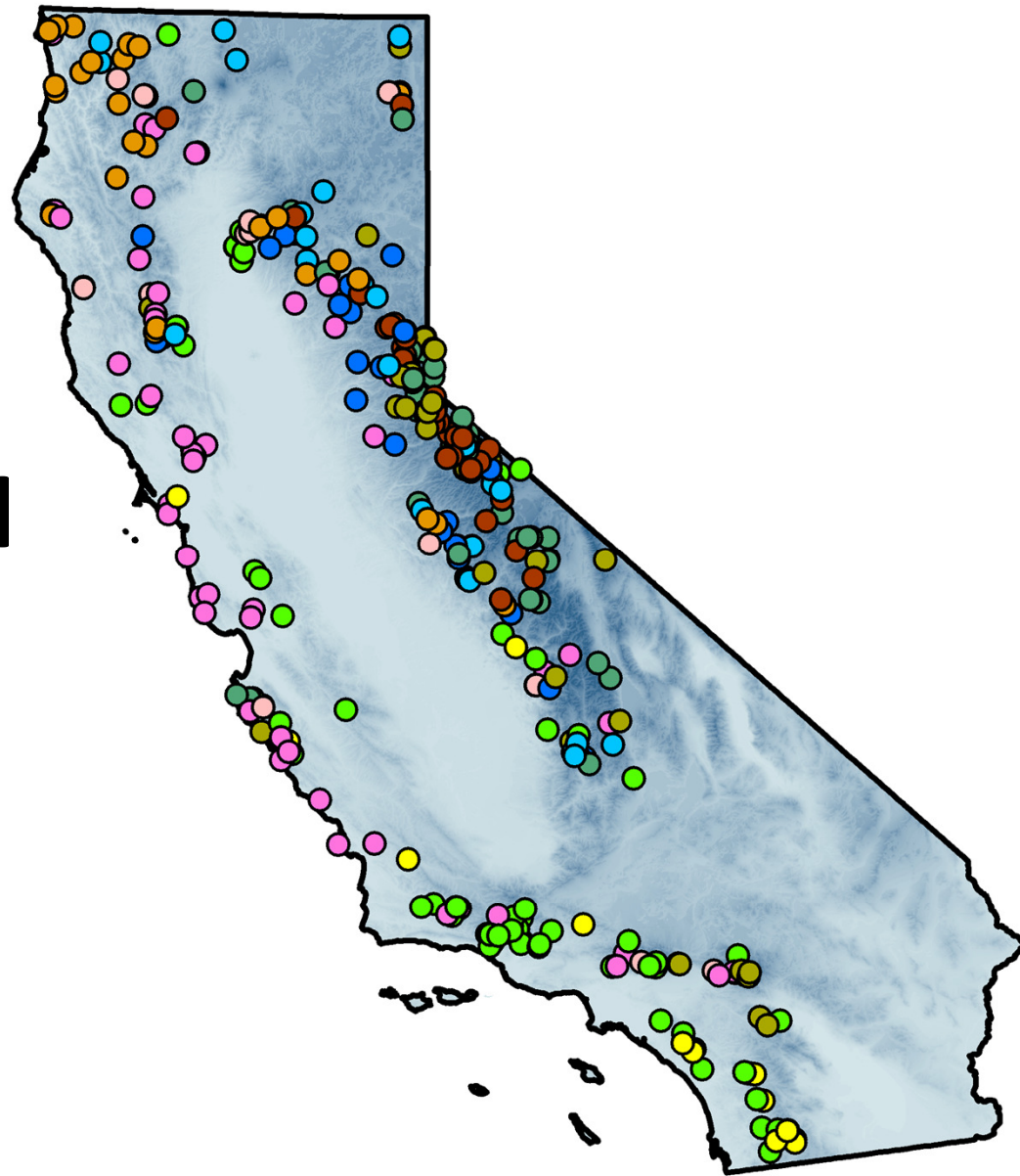
**Performance was very similar to our April O/E index**

# Cluster biological similarity

*(Bray-Curtis dissimilarity, flexible- $\beta = -0.25$ , rare taxa removed if  $< 5\%$  of sites)*



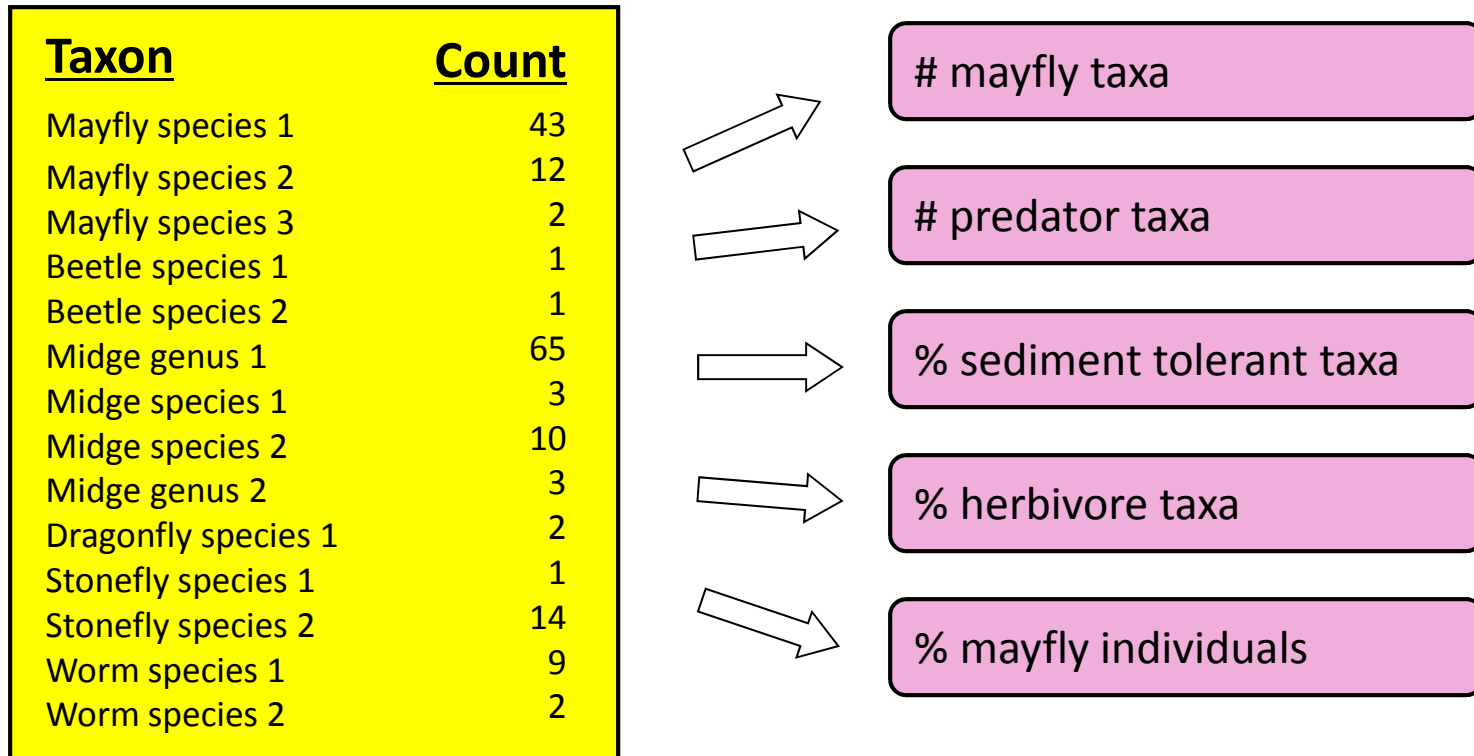
**10 biological  
clusters**





# Multi-metric Indices (MMIs)

Species list is converted into metrics representing diversity, ecosystem function, and sensitivity to stress



# Why develop an MMI?

- Science panel recommended exploring MMI
- MMIs have useful qualities
  - Measure ecological attributes other than species loss
  - Very responsive to stress
  - May work well where species-specific predictions are tricky
- **New techniques available** (*see Hawkins and Vander Laan presentation at 2011 CABW*)
  - Adds site-specific adjustments to traditional MMIs

# Building a predictive MMI (pMMI)

*follows methods of Hawkins and Vander Laan*

**Step 1.** Calculate lots of metrics at reference and stressed sites

**\*Step 2.** Create models that adjust metric values to account for major natural sources of metric variation

**Step 3.** Select metrics based on ability to discriminate reference from stressed sites

**Step 4.** Score metrics (after Cao et al. 2007) and assemble into composite pMMI



# Step 1. Calculate metrics at reference sites and stressed sites

- **Sample Information:**

- 1520 sites had “adequate” samples (i.e., >450 bugs) = 2813 samples
- 514 are reference (same definition as O/E)
- 175 are highly stressed (84 Ag, 91 Urb)
- The rest are “test”

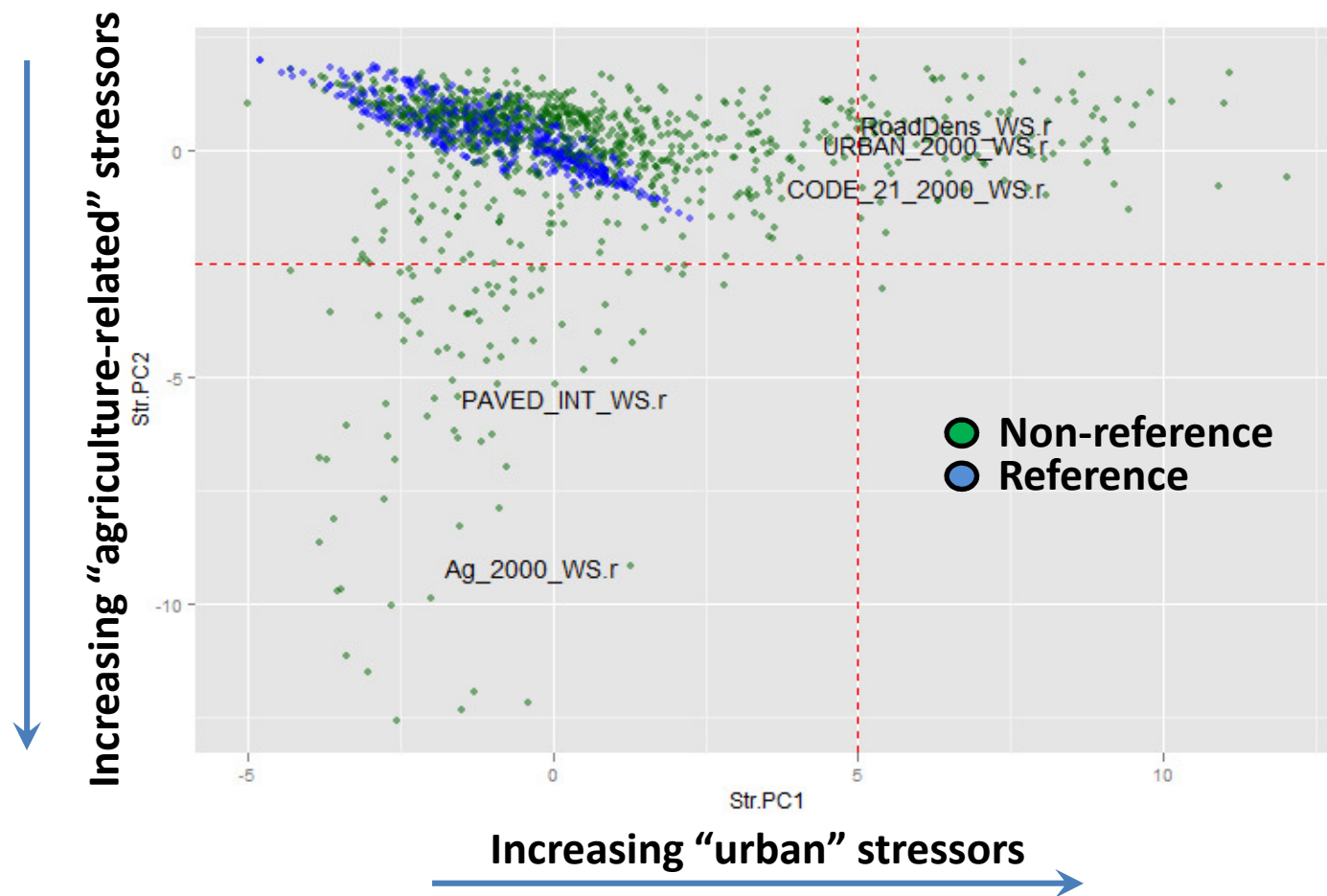
- **Calculate Metrics**

- Used SWAMP’s new bioassessment reporting module
- Subsample to 500 organisms, calculate at SAFIT Level 1 (midges to family)
- Reject samples <450 specimens

- **Use 80% for model development, 20% to validate**

# Identifying stressed sites

PCA with all GIS stressor variables (after removing effects latitude, longitude, and elevation) – stress cutoffs arbitrary



# Metrics: the usual suspects

Class	Abundance-based	# Taxa	% Taxa
Taxonomic	% EPT	EPT taxa	% EPT taxa
	[not considered]	Coleoptera taxa	% Coleoptera taxa
	[not considered]	Diptera taxa	% Diptera taxa
	% Chironomidae	[NA]	[NA]
	[not considered]	Non-insect taxa	% Non-insect taxa
	Shannon Diversity	Taxonomic richness	
FFG	% Collectors	Collector taxa	% Collector taxa
	% Predators	Predator taxa	% Predator taxa
	% Scrapers	Scraper taxa	% Scraper taxa
	% Shredders	Shredder taxa	% Shredder taxa
Tolerance	% Intolerant	Intolerant taxa	% Intolerant taxa
	% Tolerant	Tolerant taxa	% Tolerant taxa
	Weighted tolerance value		



## Step 2. Adjust metric values to account for influence of natural gradients

- Random forests models (1000 trees) allow us to predict site-specific reference expectation for each metric
- Most influential gradients (all GIS-based):

<ul style="list-style-type: none"><li>• Latitude</li><li>• Longitude</li><li>• Elevation Range</li><li>• Site Elevation</li><li>• Precipitation</li><li>• Temperature</li><li>• log Watershed Area</li></ul>	<ul style="list-style-type: none"><li>• Soil Erodability</li><li>• Soil Bulk Density</li><li>• Soil Permeability</li><li>• Hydraulic Conductivity</li></ul>	<ul style="list-style-type: none"><li>• MgO_Mean</li><li>• Surfur_Mean</li><li>• SumAve_Phos</li><li>• CaO_Mean</li><li>• Mean Phosphorus</li><li>• Mean Nitrogen</li></ul>
--	---	---

- If  $Rsq \geq 10\%$ , use metric residuals (observed – predicted). Otherwise, use raw value

## Step 3. Select most responsive metrics

- Select metrics with the best ability to discriminate reference from stressed (i.e., highest t-values – **all > t=10**)
- Avoid selecting redundant metrics
  - If  $R^2$  with any previously selected metric **> 0.5**, do not select
  - Avoid “philosophical redundancy”(e.g., EPT taxa and % EPT)

## Step 4. Score metrics and assemble into composite pMMI (*follows Cao et al. 2007*)

- **Score metrics**
  - Decreasing metrics:  $(\text{Obs} - \text{Min}) / (\text{Max} - \text{Min})$
  - Increasing metrics:  $(\text{Obs} - \text{Max}) / (\text{Min} - \text{Max})$ 
    - Max = 95<sup>th</sup> percentile of reference
    - Min = 5<sup>th</sup> percentile of stressed
- **Sum** 10 metrics and **adjust** scale to be equivalent to O/E (divide score by mean of reference)

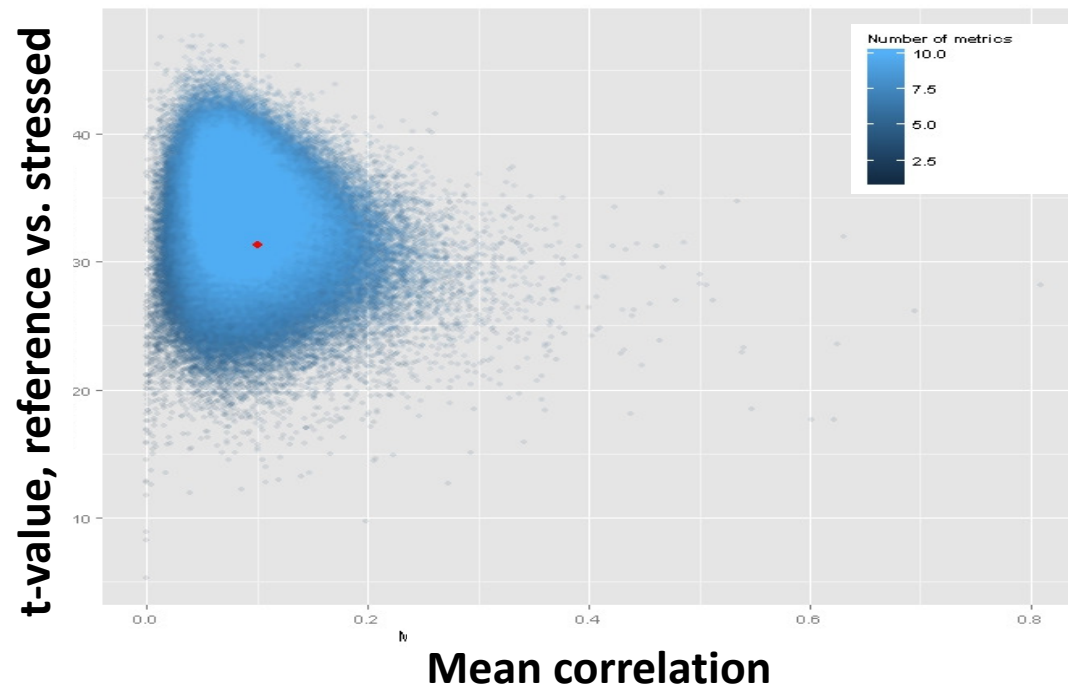
# Final Metrics

Metric	Mod v Null	% explained by RF model	t	Response
Collector taxa	Modeled	11	13.2	Decrease
Coleoptera taxa	Modeled	40	17.6	Decrease
Diptera taxa	Null	7	13.5	Decrease
Intolerant taxa	Modeled	53	32.2	Decrease
Predator taxa	Modeled	11	13.6	Decrease
Scraper taxa	Modeled	38	20.0	Decrease
Shredder taxa	Modeled	42	19.1	Decrease
% Non-Insect Taxa	Modeled	15	18.1	Increase
Shannon diversity	Modeled	16	10.7	Decrease
Tolerance value	Modeled	32	12.4	Increase

# Evaluated multiple MMIs

All subsets of 30 metrics

(~100,000 MMIs; 10 metrics max, no redundancy)

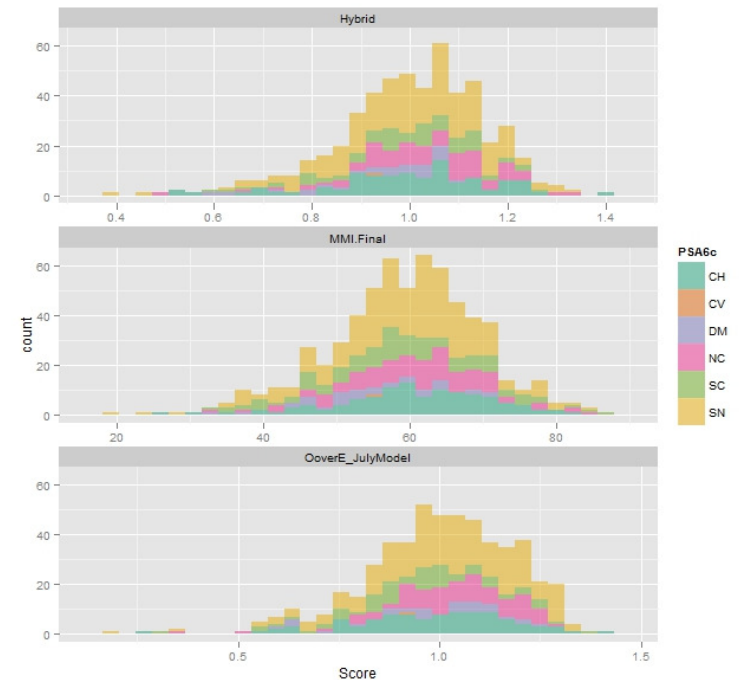


- Nearly all MMIs discriminate (reflects pre-screening of metrics?)
- More metrics = convergence to central tendency, better validation
- Thousands of other MMIs are probably just as good as ours



# Comparing Performance of 3 Scoring Tools

1. Species Loss Index (O/E)
2. Ecological Structure Index (pMMI)
3. Combined Index (“hybrid”)



**Created a common validation set for performance measures  
so we're comparing apples to apples**

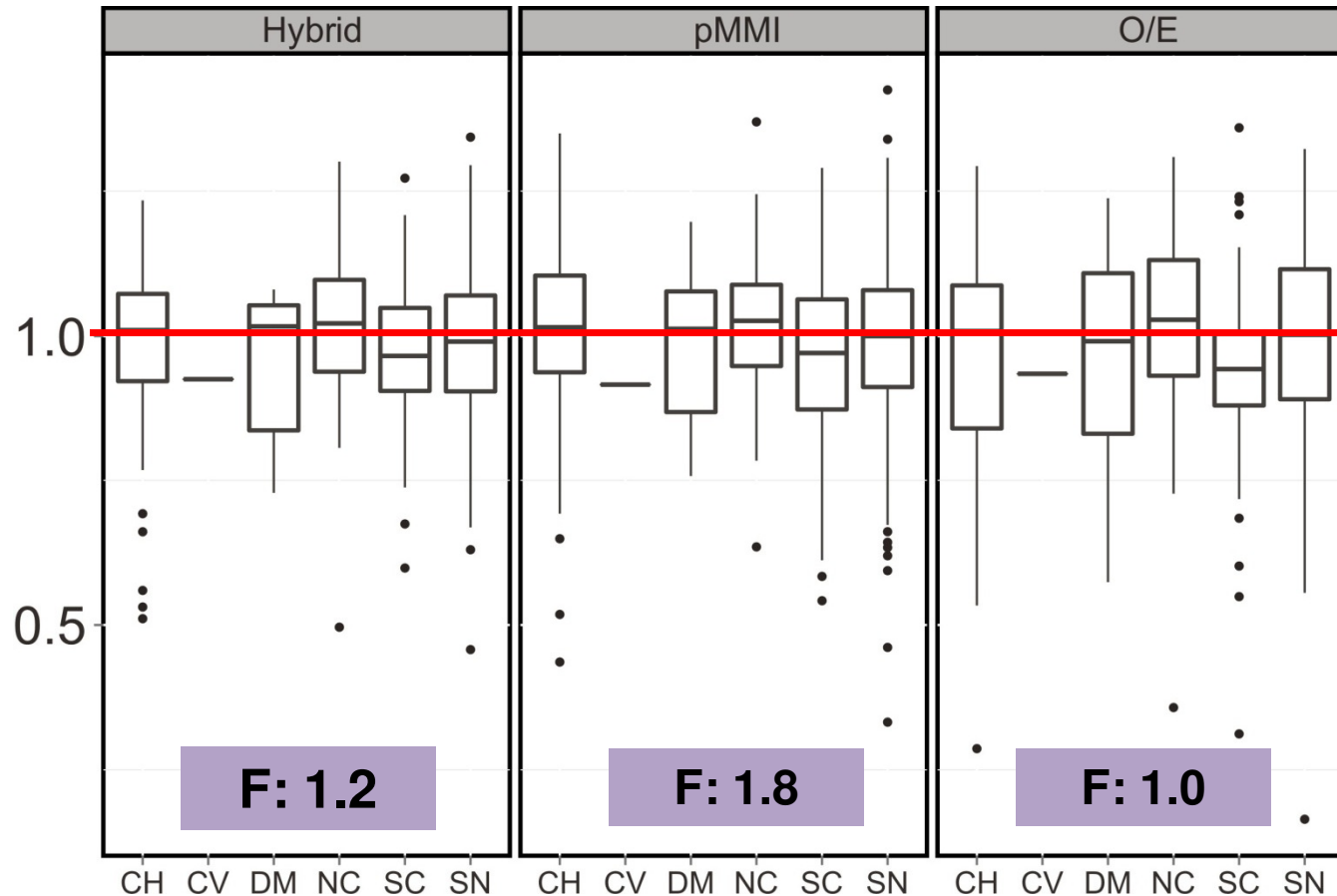
# Measuring Performance

All evaluations used a common dataset

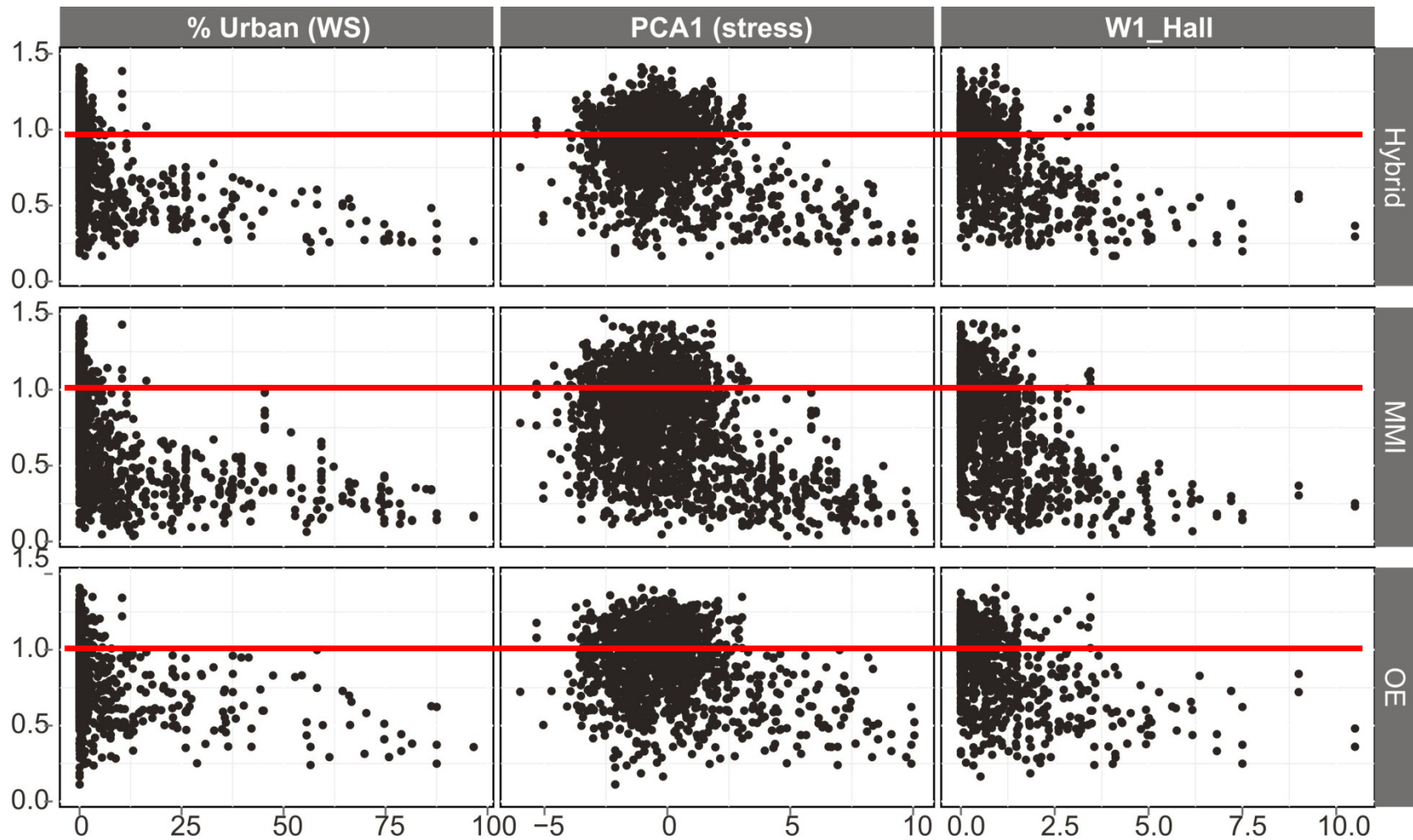
Class	Property	Measure	O/E	pMMI	Hybrid
Precision	Variance of reference sites	SD	0.19	0.15	0.14
Sensitivity/ Responsiveness	Discrimination	t-value	9.5	17.6	15.3
	Variance explained by stress	Random forest model	25%	56%	49%
Accuracy/ Bias	Variance explained by natural gradients (ref sites)	Random forest model	-7%	-9%	-8%
	Difference among PSA regions (ref sites)	ANOVA	1.0 (ns)	1.8 (ns)	1.2 (ns)
Replicability	Within-site variability	Mean within-site SD	0.10	0.10	0.08

# Statewide Consistency

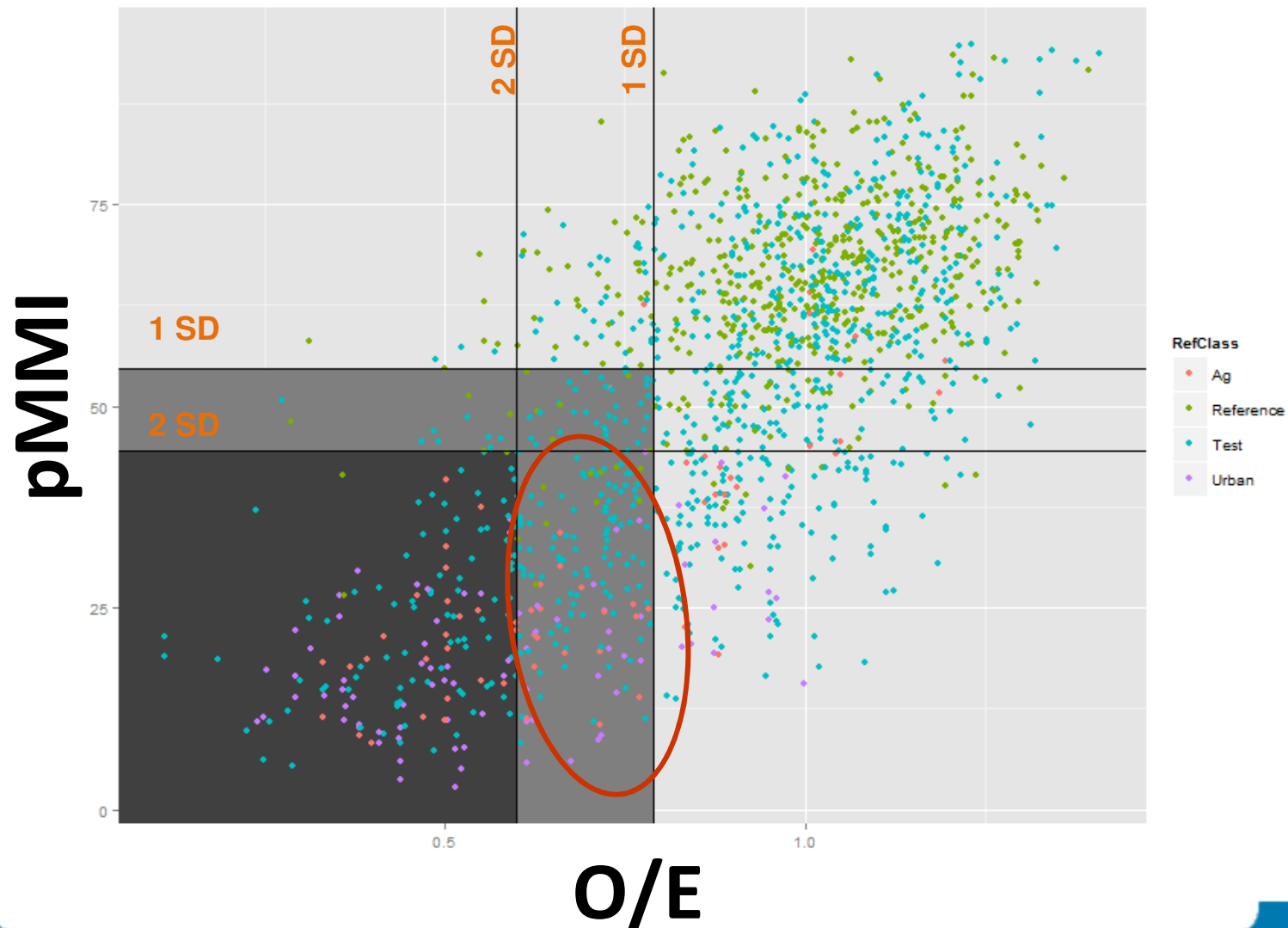
Distribution of reference scores by PSA region



# Responsiveness to stress

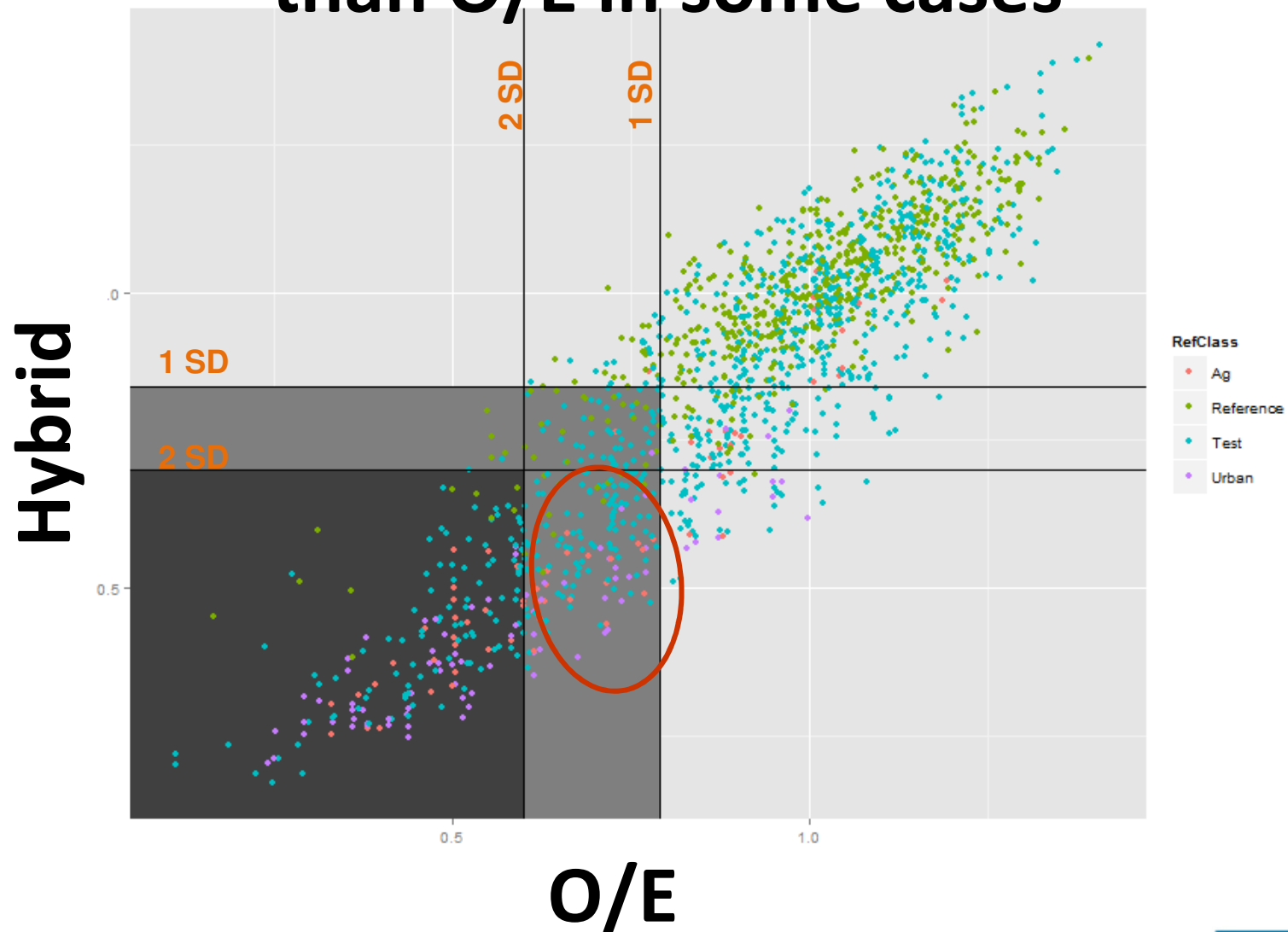


**pMMI and O/E have general agreement,  
but tell us somewhat different things**

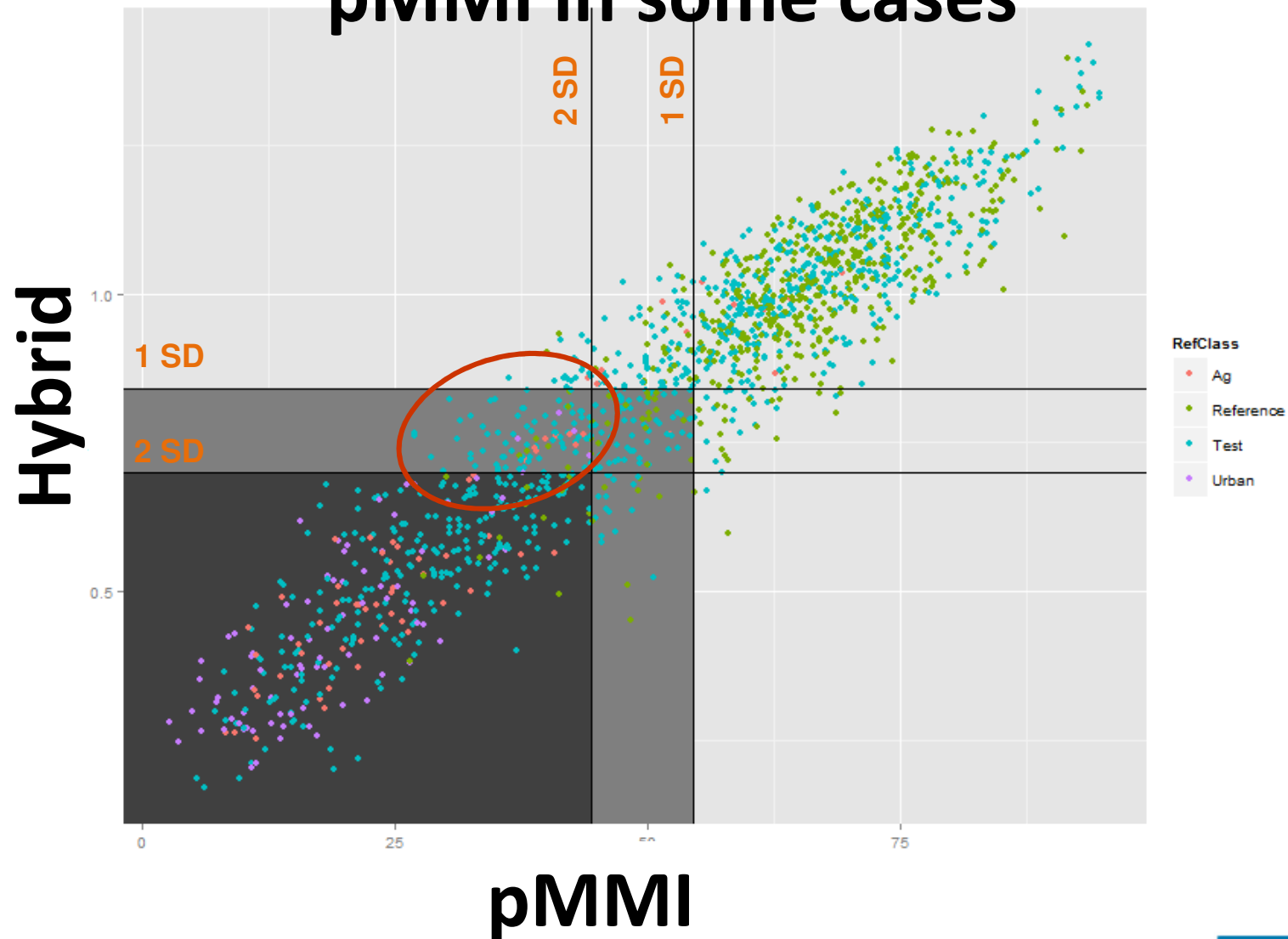




# Hybrid more likely to find impairment than O/E in some cases



# Hybrid less likely to find impairment than pMMI in some cases



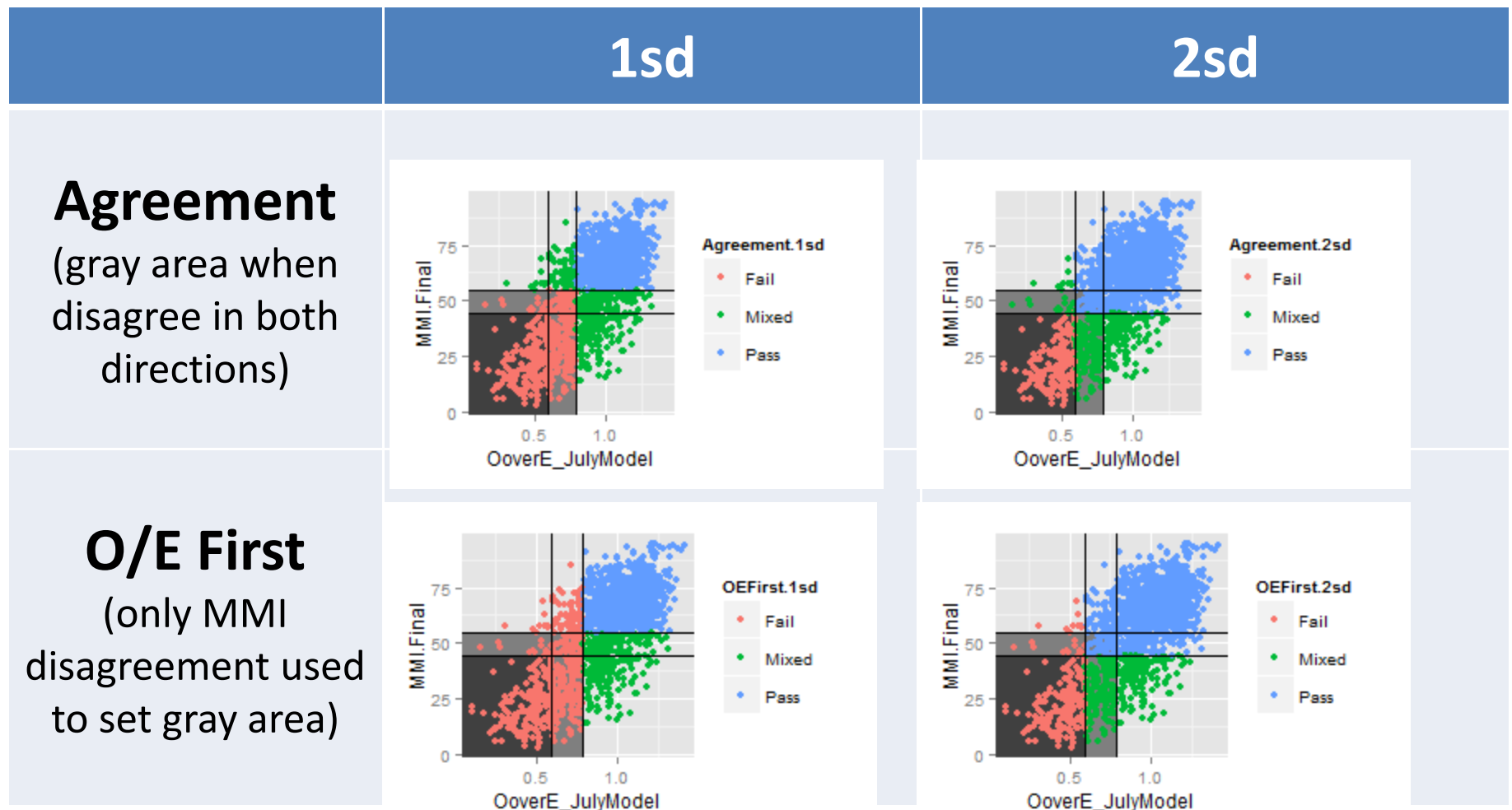
## **Both pMMI and O/E have desirable qualities**

- pMMI is precise and very responsive to stress (but it was designed to be)
- % species loss is an intuitive, meaningful measure of condition
- Both are accurate and applicable throughout state
- Potential for complementarity is great -- we explored a few options (see Science Panel)

# Options for using 2 indices

- Hybrid
  - Equal
  - Unequal weight
- Agreement/ Disagreement
- Use one to verify the other

# Multi-index Approaches





# We recommend an equal-weight combined index

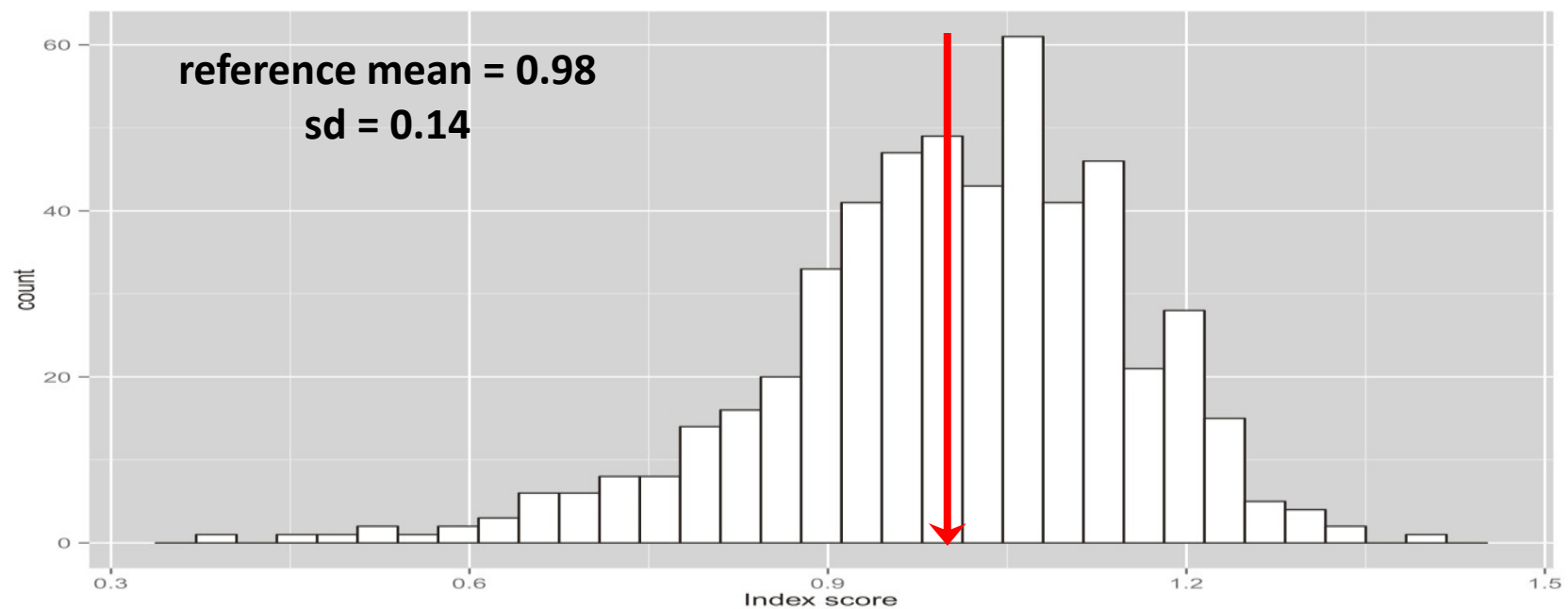
- Retains some of the better qualities of both indices, tempers weaknesses
- Retains the precision and high sensitivity of the MMI and the independence of the species loss data
- Can be disaggregated into component MMI and O/E
  - Don't lose information by combining
  - Reference expectations for all components are available
- No objective *a priori* reason to weight
- Implementation is easier with a single score

# California Stream Condition Index (CSCI)

Part A: Ecological Structure Component (pMMI)

Part B: Taxonomic Loss Component (O/E)

**CSCI is a simple average of the two scores**



# Options for setting thresholds

- **Statistical criteria**
  - Standard deviation
  - %-ile of reference distribution
- **Ecological criteria**
  - Acceptable species loss or change in community structure



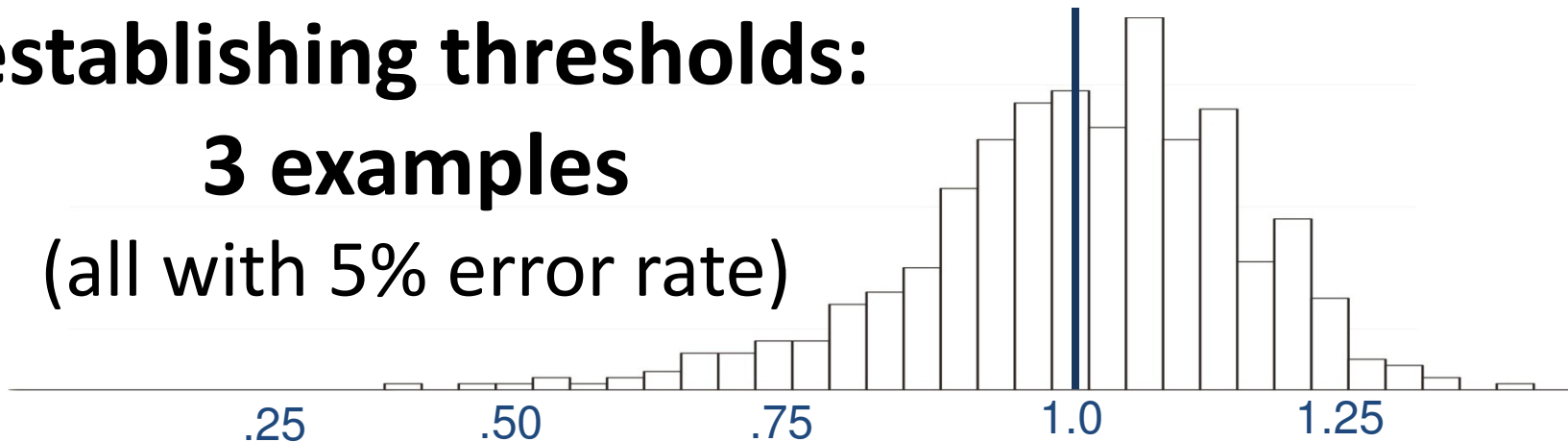
# **We recommend statistically defined thresholds with a gray area**

- Widely accepted practice with broad acceptance
- Ecological benchmarks are appealing biologically, but we have limited basis for setting these objectively
- Gray area is helpful way to express uncertainty in whether a sample reflects site condition

# Statistical approaches to establishing thresholds:

## 3 examples

(all with 5% error rate)



95% and 85% confidence that site is not equivalent to reference



95% confidence that the 95% threshold is where we think it is



Use within-site error rate to establish uncertainty around threshold

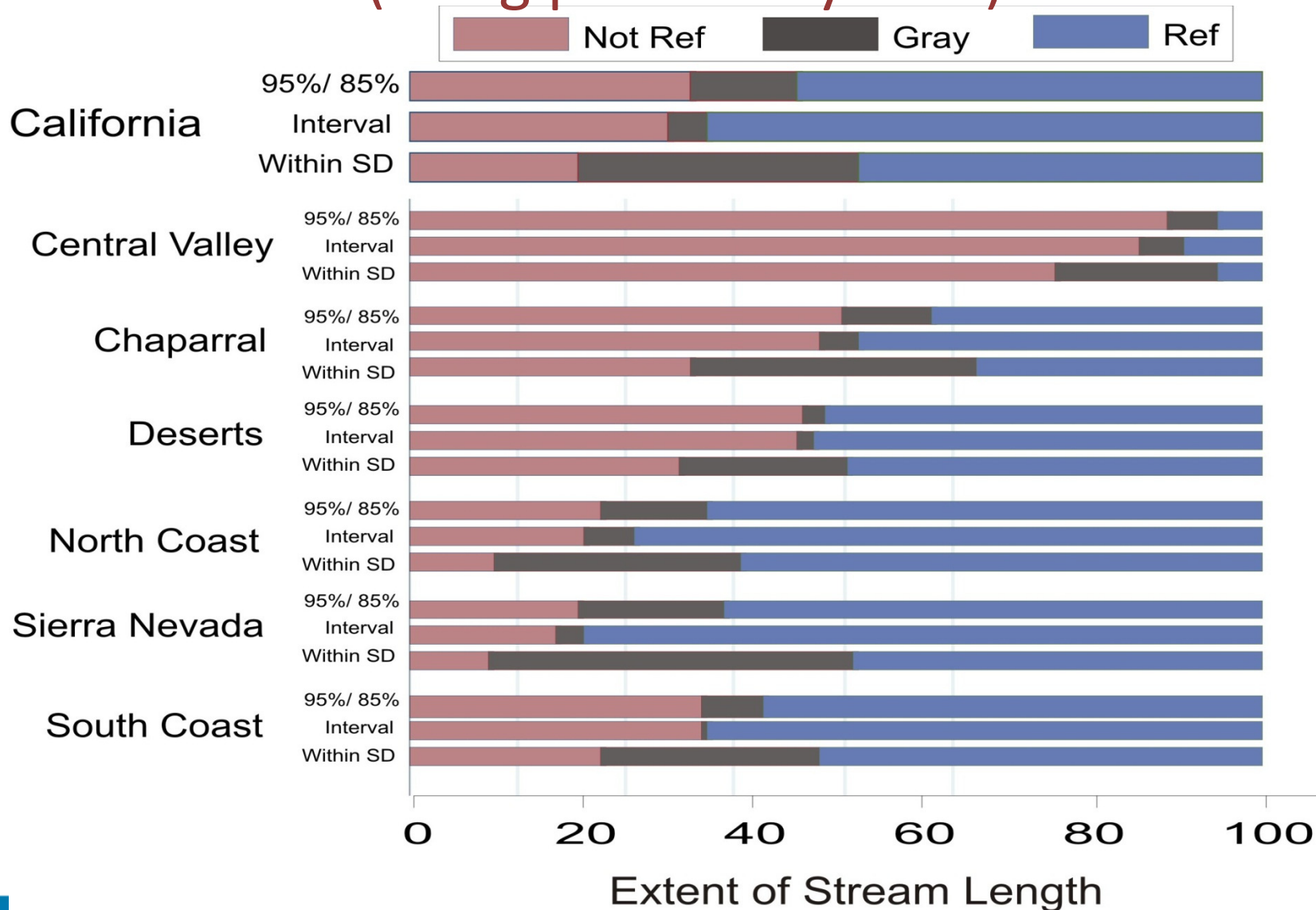


# Different approaches for multiple samples (i.e., increasing certainty about site condition)

- Formal t-test vs. threshold
  - **Pass** if site mean  $>$  threshold; **Fail** if site mean  $<$  threshold
  - **Gray**: mean  $\sim$  threshold
    - Different responses given power of the test
      - Low power: More sampling
      - High power: Apply strict threshold comparison (no gray zone)
- What about Type II error?
  - Compare test distribution to reference distribution?
  - Set alpha at 0.10 or higher?
- Other ideas?

# Extent of stream length by region

(using probability data)



# Questions about thresholds

Are there other options we should consider for guarding against Type I error (false positives)?

Can you suggest objective ways to protect against Type II error (false negatives)? Is there a way to incorporate a “safety factor”?

Do you favor one of the approaches for bounding a gray area?

Can you recommend strategies for dealing with multiple data points?

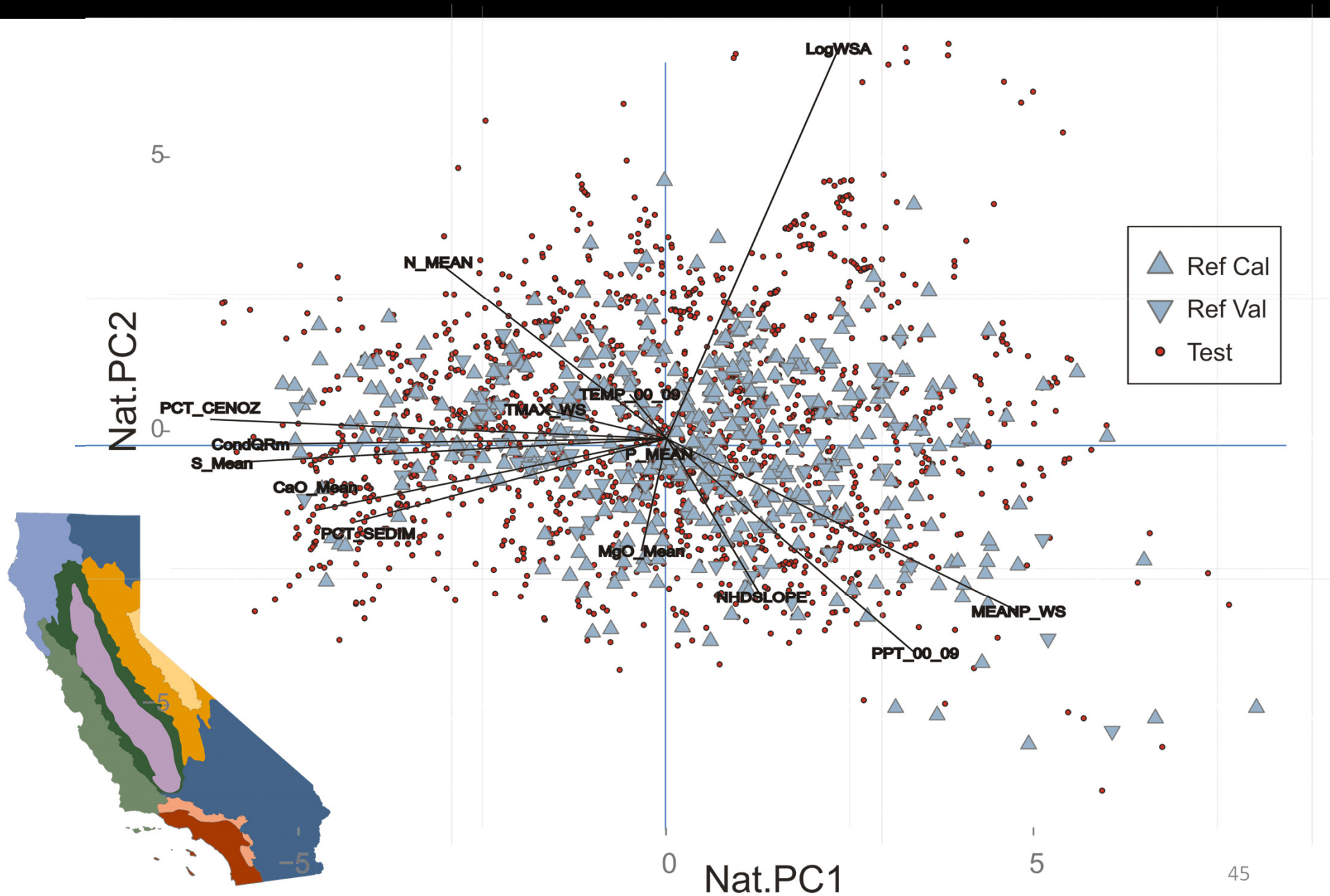
# What's next? (Part I):

## Quantify applicability of tool

**Goal:** develop an objective means for determining whether a test site can be appropriately scored(i.e., “is a test site within the “experience” of the model)

- Develop a multivariate applicability test (e.g., Mahalanobis distance)?
- Univariate tests?
- Other ideas?
- How do we define a criterion of acceptance?
- Could be a good way to establish exceptions for truly unique environmental settings.

# Multivariate view of natural diversity



# What's next? (Part II)

## Automation and Documentation

### Automate calculations

- Package GIS layers
- Make standard calculation and reporting tools available

### Document, document, document

- Journal articles
- Website 101 and FAQ
- Website appendices



# Questions for the panel (Part I)

- Are our scoring tools ready to support implementation?
- Are there other factors we should consider before finalizing our scoring tool recommendations?
  - Combination index versus other options
  - Inclusion of a grey area or not
  - Balancing Type I and II errors
- Gray area options
  - Should we explicitly deal with multiple data points in our gray area approach?

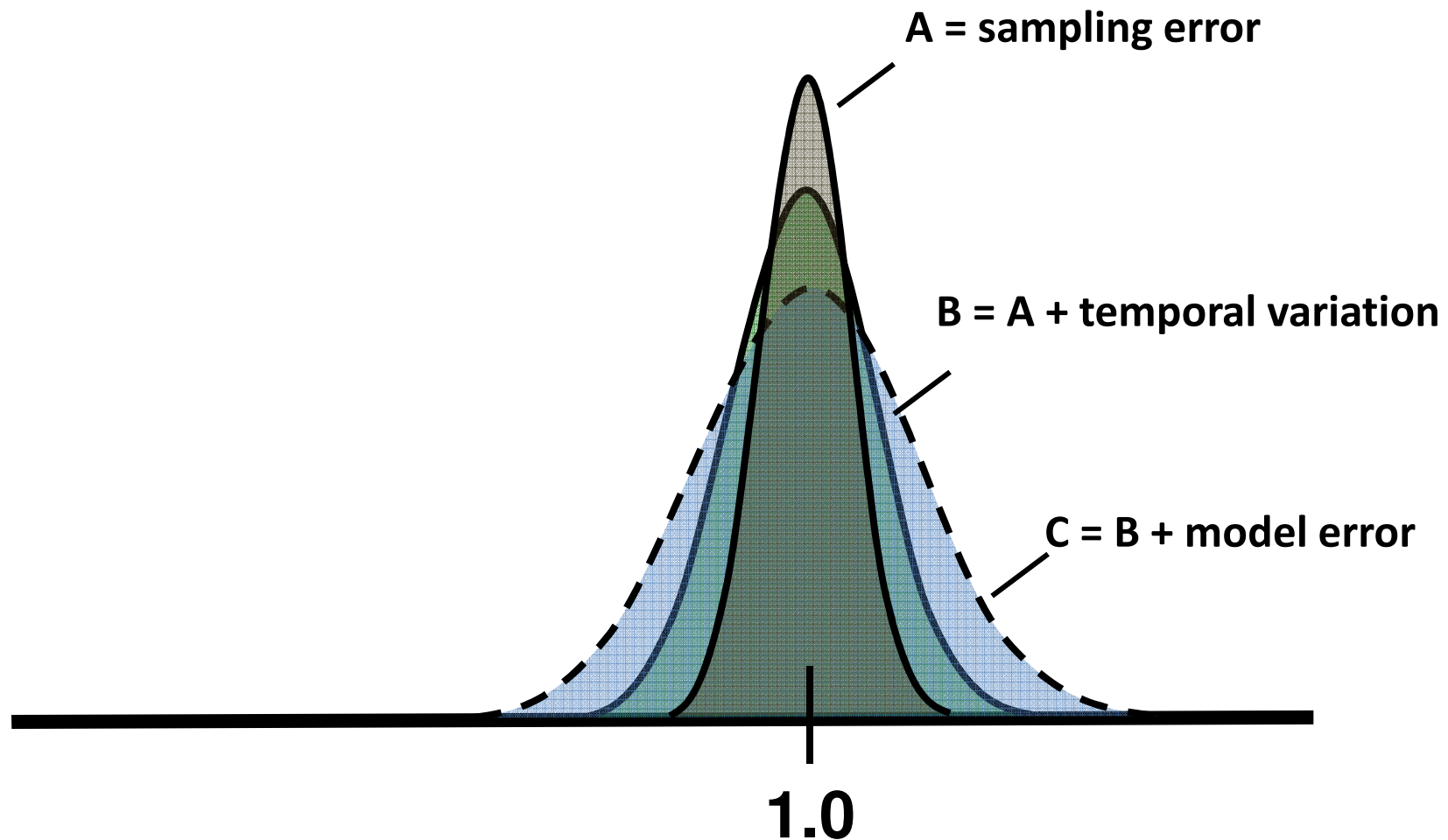


# Questions for the panel (Part II)

- Recommendations for exploring and quantifying limits of tool?
- Recommendations for automation?
- Recommendations for documentation?



# Sources of variation in index scores



(after Hawkins et al. 2010)